

Figure S1. SARS-CoV-2 S protein induces inflammatory cytokines in macrophages. (A) THP-1 cells were stimulated with S2 protein. 4h post stimulation, culture supernatants were collected, centrifuged, and analyzed for IL-6, IL-1 β , and TNF α by ELISA. (B) THP-1 cells were stimulated with PolyI:C (1 μ g/ml). The expression of interferons was measured by real-time PCR. (C) THP1 cells were stimulated with native S2 or heat-denatured S2 (500 ng/ml). The induction of IL6, IL1b, and TNFa was measured by real-time PCR at 4h post stimulation. Data represent mean \pm SD (n=3); **p < 0.001 by unpaired Student's t test. Experiments were repeated two times and data of a representative experiment is presented.

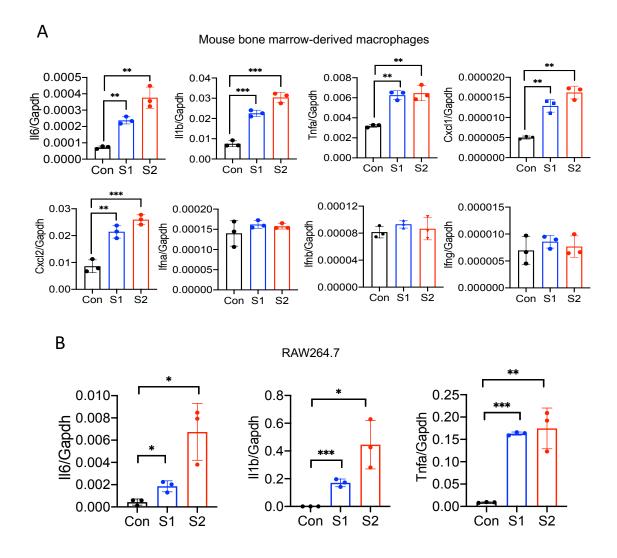


Figure S2. Mouse macrophages are stimulated by SARS-CoV-2 S protein. (A) Bone marrow-derived macrophages from WT mice were stimulated with S1 and S2 proteins (500 ng/ml) for 4 h. The expression of *Il6*, *Il1b*, *Tnfa*, *Cxcl1*, *Cxcl2*, *Ifna*, *Ifnb*, *Ifng* was measured by real-time PCR. (B) RAW264.7 murine macrophage cells were stimulated with S1 or S2 (500 ng/ml) for 4h. The expression of *Il6*, *Il1b*, and *Tnfa* was measured by real-time PCR. Data represent mean \pm SD (n=3); *p < 0.05, **p < 0.001, ***p < 0.0001 by unpaired Student's t test. Experiments were repeated two times and data of a representative experiment is presented.

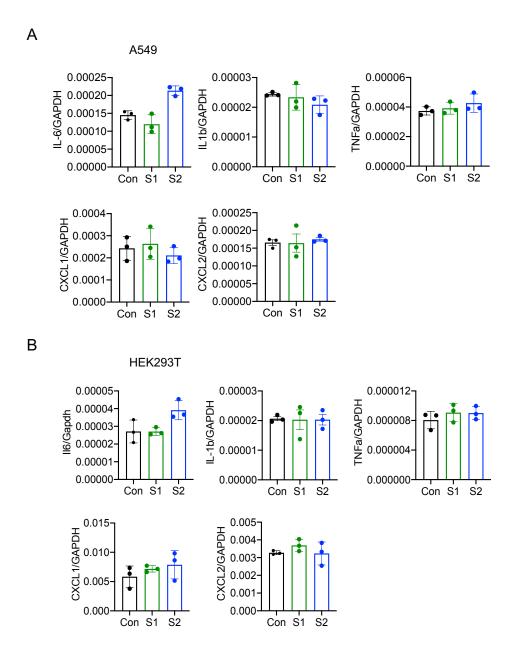


Figure S3. Epithelial cells do not respond to SARS-CoV-2 S protein acutely. (A-B) A549 or HEK293T cells were incubated with SARS-CoV-2 S1 or S2 (500 ng/ml) proteins for 4 h. The expression of inflammatory cytokines and chemokines was measured by real-time PCR. Data represent mean \pm SD (n=3). Experiments were repeated two times and data of a representative experiment is presented.

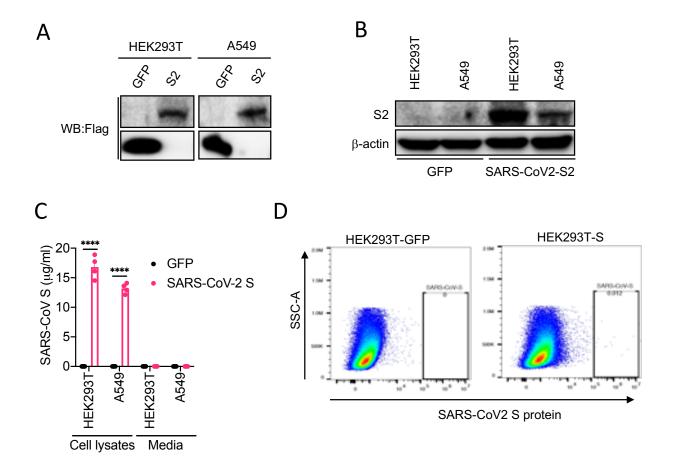


Figure S4. Expression analysis of SARS-CoV-2 S protein in cultured cells. HEK293T or A549 cells were transfected with plasmids containing flag-tagged S or green fluorescent protein (GFP). (A-B) 48h post transfection, cell lysates were collected and the expression of S was measured by western blot analysis of Flag (A) and S (B) using respective antibodies. (C) Cell culture supernatants and cell lysates were collected at 48h after transfection, and analyzed for S protein by ELISA. (D) The expression of S on the cell surface of HEK293T-S cells was measured by flow cytometry following surface staining of S proteins. Data represent mean \pm SD; ****p<0.0001 by unpaired Student's t test. Experiments were repeated two times and data of a representative experiment is presented.

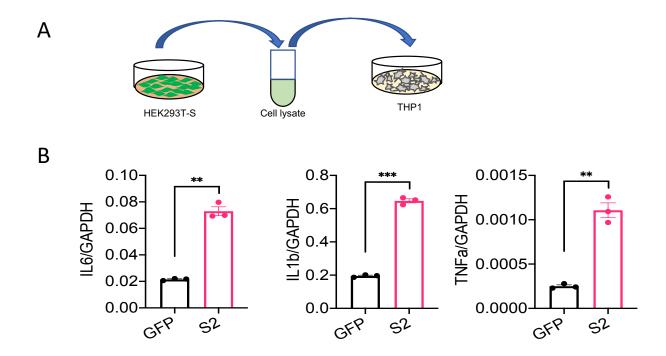


Figure S5. Cell lysate of HEK293T cells expressing S protein activates macrophages and induced inflammatory cytokines. (A) S or GFP (control) proteins were overexpressed in HEK293T cells. Cells were sonicated, and cell lysate supernatants were collected. THP-1 cells were incubated with these cell lysate supernatants for 4 h. (B) The expression of *IL6*, *IL1b*, and *TNFa* in THP cells was measured by real-time qPCR. Data represent mean \pm SD (n=3); **p < 0.001, ***p < 0.0001 by unpaired Student's t test. Experiments were repeated two times and data of a representative experiment is presented.

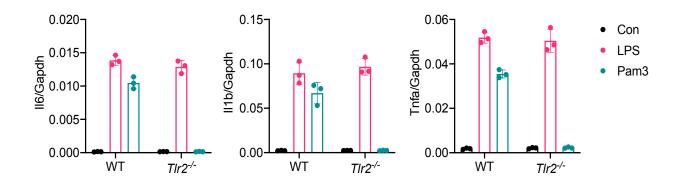


Figure S6. Macrophages of $Tlr2^{-/-}$ mice are defective in sensing TLR2 ligand Pam3CSK4. BMDMs from WT and $Tlr2^{-/-}$ mice were stimulated with LPS (1µg/l) or Pam3CSK4 (1µg/ml) for 4 h. The expression of inflammatory cytokines was measured by real-time PCR. Data represent mean \pm SD (n=3).